

**AMENDMENTS TO THE CLAIMS**

Claims 8 and 21-23 are amended. A complete listing of the claims is provided below.

**1-7 (Cancelled)**

**8. (Currently amended)** A method for in vitro screening for a peptide capable of altering the phenotype of a cell, said method comprising the steps:

- a) introducing a molecular library of retroviral vectors comprising randomized nucleic acids into a plurality of cells to provide for expression of a plurality of test peptides each comprising a randomized amino acid sequence and a glycine N-terminal to said randomized amino acid sequence,
- b) screening said plurality of cells for a cell exhibiting an altered phenotype due to an interaction between a test peptide and a cellular component native endogenous to said cell; and
- c) identifying said peptide capable of altering the phenotype of said cell.

**9. (Previously presented)** A method according to claim 8 wherein said identifying comprises:

- i) isolating said cell exhibiting an altered phenotype.

**10. (Previously presented)** A method according to claim 9 wherein said identifying further comprises:

- ii) sequencing said nucleic acid encoding said peptide capable of altering the phenotype of said cell.

**11. (Previously presented)** A method according to claim 8 wherein said nucleic acids further encode a presentation sequence capable of presenting said test peptides in a conformationally restricted form.

**12. (Previously presented)** A method according to claim 8 wherein said cells are mammalian cells.

13. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^4$  different nucleic acids.

14. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^5$  different nucleic acids.

15. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^6$  different nucleic acids.

16. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^7$  different nucleic acids.

17. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^8$  different nucleic acids.

18. **(Previously presented)** A method according to claim 8 wherein said library comprises at least  $10^9$  different nucleic acids.

19. **(Previously presented)** A method according to claim 8 wherein each of said nucleic acids is linked to nucleic acid encoding at least one fusion partner.

20. **(Previously presented)** A method according to claim 19 wherein said fusion partner comprises a nuclear localization signal sequence.

21. **(Currently amended)** A method for in vitro screening for a peptide capable of altering the phenotype of a cell, said method comprising the steps:

a) introducing a molecular library of retroviral vectors comprising randomized nucleic acids into a plurality of cells to provide for expression of a plurality of test peptides each comprising a randomized amino acid sequence;

b) screening said plurality of cells for a cell exhibiting an altered cell growth phenotype, wherein said altered phenotype is due to an interaction between a test peptide and a cellular component native endogenous to said cell; and

c) identifying said peptide capable of altering the cell growth phenotype of said cell.

22. **(Currently amended)** A method for in vitro screening for a peptide capable of altering the phenotype of a cell, said method comprising the steps:

a) introducing a molecular library of retroviral vectors comprising randomized candidate nucleic acids into a plurality of cells to provide for expression of a plurality of test peptides each comprising a randomized amino acid sequence;

b) screening said plurality of cells for a cell exhibiting an altered cell death phenotype, wherein said altered phenotype is due to an interaction between a test peptide and a cellular component native endogenous to said cell; and

c) identifying said peptide capable of altering the cell death phenotype of said cell.

23. **(Currently amended)** A method for in vitro screening for a peptide capable of altering the phenotype of a cell, said method comprising the steps:

a) introducing a molecular library of retroviral vectors comprising randomized candidate nucleic acids into a plurality of cells to provide for expression of a plurality of test peptides each comprising a randomized amino acid sequence;

b) screening said plurality of cells for a cell exhibiting a change in expression of a cellular differentiation marker, wherein said change in expression is due to an interaction between a test peptide and a cellular component native endogenous to said cell; and

c) identifying said peptide capable of changing expression of a cellular differentiation marker of said cell.

24. **(Previously presented)** The method according to claim 23, wherein said cellular differentiation markers are characteristic of T-cell activation.

25. **(Previously presented)** The method according to claim 23, wherein said cellular differentiation markers are characteristic of B-cell activation.